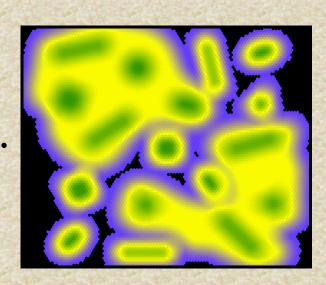
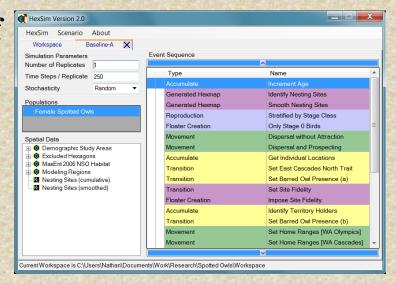


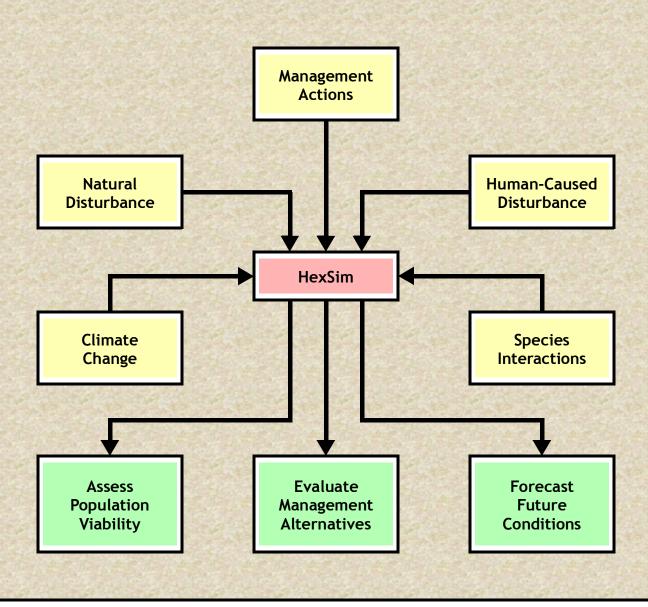
What is HexSim?

- → It is a computer simulation model.
- → It was designed for evaluating wildlife population responses to human activities.
- → It balances generality and flexibility with parsimony and ease of use.
- → It can be used with a large range of places, problems, and questions.





Using HexSim in Ecological Research



Significant Challenges

- → Landscapes. They are dynamic; structure matters; features change with life history
- → Populations. They have complex, diverse life histories, and can interact
- **→** Disturbance. Can vary in space and time; there can be multiple; they often interact
- → Methodology. Must be defensible and usable, plus have value to decision-makers

How is HexSim Different?

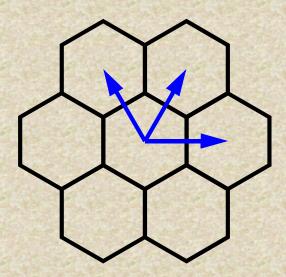
- → It has a wide range of potential applications.
- → It contains no simplifying assumptions about the biology or ecology of the study systems
- → Every individual can have unique properties that change throughout their lifetimes
- → Can simulate population interactions, stressor interactions, landscape genetics, and more
- → Modern and easy to use, with graphical user interfaces (GUI) for every model component

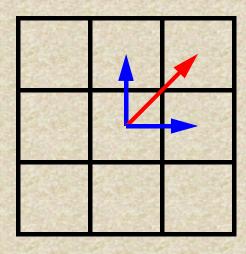


- → Spatially-Explicit and Individual-Based
- **→** Landscapes can Change Continuously
- **→** No Built-In Assumptions or Rules
- **→** Multi-Stressor with Interactions
- **→** Multi-Population with Interactions
- **→** Females-only or 2-Sex Simulations
- → Life History Events Stratified by Traits

Why Hexagons?

- → They provide a space-filling tesselation
- → Each of a hexagon's neighbors is the same distance away.





Model Inputs

- → Spatial Data. Can be real or fabricated, one or multiple layers, static or time series...
- → Life History Data. Can be real or fabricated or a hybrid. Data limits model complexity...
- → Disturbance Regimes. Spatial, temporal, simple, complex, local, regional, etc...
- **→** Stochasticity. Demographic, environmental, life stage-specific, spatially-distributed, etc...

Model Outputs

- → Census Data. Chronological records of userdefined population metrics.
- → Tabular Reports. CSV files detailing observed vital rates, movements, interactions, etc.
- **→** Map-Based Reports. Map files illustrating population performance and interactions.
- → Videos. Movies showing movement, resource acquisition, occupancy by trait class, etc.

Life History Events

- → Survival
- → Reproduction
- **→** Movement
- **→** HexMap Generation
- **→** Species Interaction
- **→** Species Introduction
- **→** Mutation
- → And so on...

Trait Types

- **→** Probabilistic Traits
- **→** Accumulated Traits
- **→** Genetic Traits

Survival ⇒ Combinations ↓	Mean Rate		
Juvenile	0.7		
Subadult	0.8		
Adult	0.9		

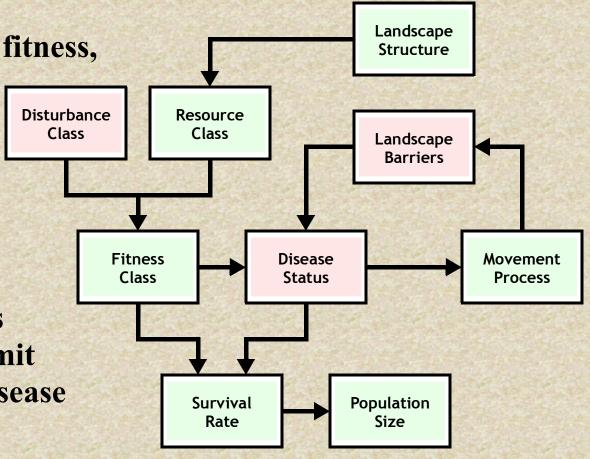
	Births ⇒ Combinations ∜	0	1	2	3	4	5	6	Expected Value
٠	Female, Juvenile	1	0	0	0	0	0	0	0
	Female, Adult	0	0.1	0.1	0.2	0.3	0.2	0.1	3.7
	Male, Juvenile	1	0	0	0	0	0	0	0
	Male, Adult	1	0	0	0	0	0	0	0

A Hypothetical HexSim Scenario of Moderate-Complexity

Disturbance affects fitness, which in turn impacts disease status, survival,

and reproduction

Movement barriers affect survival rates because they can limit the spread of the disease



HexSim Genetics

- → Each individual is assigned a genotype
- → Populations can have any number of loci
- → Each locus can have any number of alleles
- → Inheritance can be from mother, father, or from both parents (per locus)
- → User-defined initial conditions, include spatial stratification of alleles

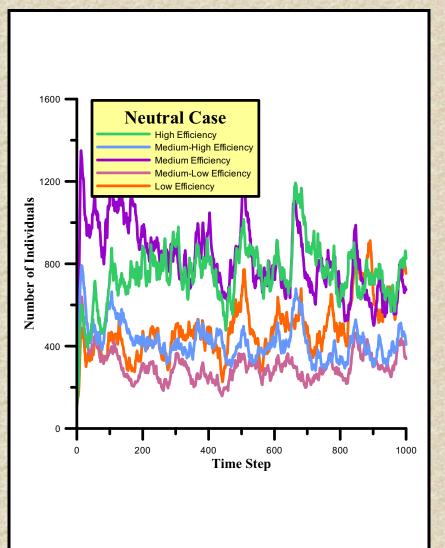
HexSim Genetics (cont.)

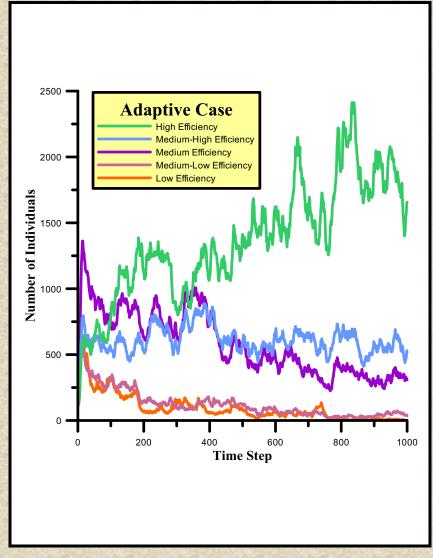
- → Mutation events my be influenced by non-heritable traits (e.g. exposure)
- → Heritable traits may be neutral or adaptive
- → Heritable and other traits may be combined to influence life history events
- → Map-distances may be used to simulate chromosome crossover

Example: Predators & Prey

- **→** Two interacting populations
- → Predators & prey use different mating schemes
- > Prey live in colonies, predators do not
- Predator males disperse towards prey Predator females disperse towards males
- → Predator capture efficiency is controlled through a heritable trait.
- → Capture efficiency influences reproduction through a resource acquisition trait
- **→** Mutation alters capture efficiency trait

Allele Frequencies





Some PATCH / HexSim Applications

- → Ord's Kangaroo Rats
- **→** Spotted Owls
- **→** Kit Foxes
- **→** Lyme Disease
- **→** Pileated Woodpeckers
- **→** Desert Tortoise
- **→** Black-capped Vireos / Cowbirds
- → Elk
- **→** Wolves
- → Fishers
- → Martin

Ord's Kangaroo Rats (Alberta)



STATE OF WASHINGTON

September 2004

Feasibility Assessment for Reintroducing Fishers to Washington



by Jeffrey C. Lewis and Gerald E. Hayes







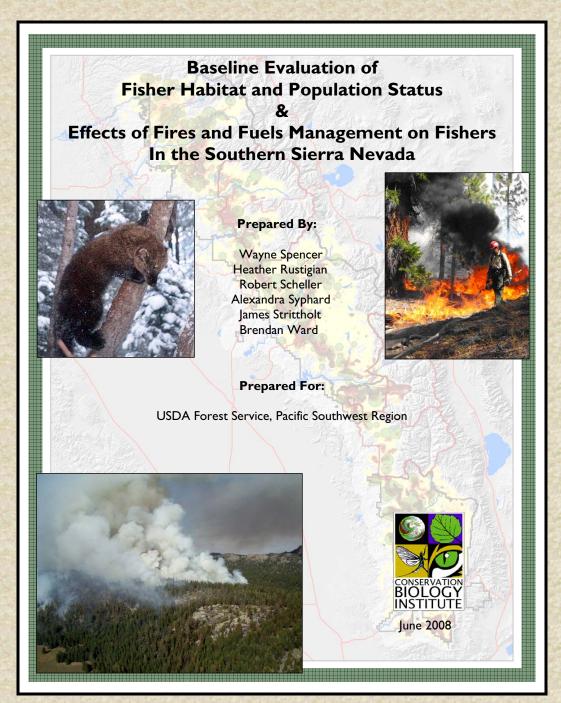
Fisher Reintroduction

Table 13. Median number of female fishers predicted by the PATCH model to be supported on potential reintroduction areas in the Olympic Peninsula, Northwestern Cascades, and Southwestern Cascades. Values were derived from 20 replicate simulations started with 30, 60 and 100 female fishers: male presence is assumed in the model.

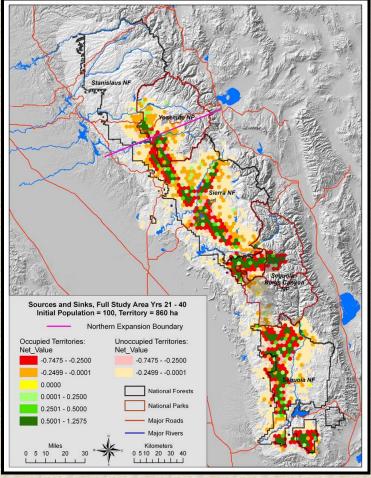
isners; male presence is a	issumed in the model.				
		Median number of female fishers supported			
Simulation specifications		Olympic	Southwestern	Northwestern	
Leslie matrices used ¹	Simulation length	Peninsula	Cascades	Cascades	
	Reintroduced				
	20 years	82.5	36	25	
Single Mean	40 years	94	33.5	19.5	
	20 years	81.5	35	17	
Six Random	40 years	84.5	29.5	17	
	60 Females R	eintroduced			
	20 years	93.5	48	26.5	
Single Mean	40 years	92	36.5	21	
	20 years	90.5	49.5	27.5	
Six Random	40 years	87.5	25	21	
100 Females Reintroduced					
	20 years	98.5	59.5	31	
Single Mean	40 years	96	43.5	20	
	20 years	102	54.5	30.5	
Six Random	40 years	87	44.5	23.5	
100 Females F	Reintroduced, additional	specifications	for sensitivity tes	ting ²	
Single Mean, 25 km	20 years	101.5	57.5	32	
maximum dispersal	40 years	97.5	48	24	
Single Mean, 75 km	20 years	98.5	55	30	
maximum dispersal	40 years	96	40	22.5	
Single Mean, low habitat	20 years	50.5	6	4.5	
scores	40 years	51.5	0	0	

Two matrix scenarios were used in simulations. The single mean simulations were run with 1 Leslie matrix with mean values for survival and fecundity. The six random matrix simulations used four matrices of mean survival and fecundity values, one matrix with low values, and one matrix with high values; one of these six matrices was chosen at random each year of a simulation to incorporate environmental stochasticity.

² Three alternative simulations were run to test the sensitivity of the model to: a smaller maximum dispersal distance of 25 km, a large maximum dispersal distance of 75 km, and lower habitat scores for suboptimal habitats.



Fisher Dynamics in the Sierra Nevada



U.S. Fish & Wildlife Service

2010 Draft Revised Recovery Plan for the Northern Spotted Owl (Strix occidentalis caurina)



Spotted Owl Recovery

Modeling Regions

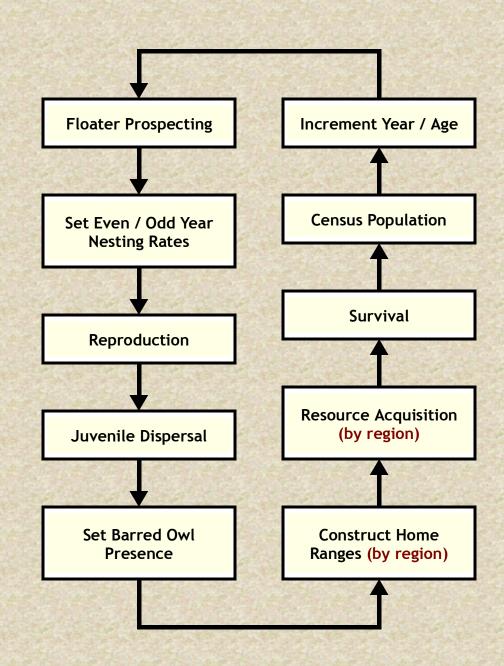
CODE	DESCRIPTION
NCO	North Coast and Olympic
ORC	Oregon Coast
ECS	Eastern Cascades - South
ECN	Eastern Cascades - North
WCN	Western Cascades - North
wcc	Western Cascades - Central
wcs	Western Cascades - South
KLE	Klamath-Siskiyou - East
KLW	Klamath-Siskiyou - West
ICC	Interior California Coast
RDC	Redwood Coast

No warranty is made by the U.S. Fish and Wildlife Service as to the accuracy, reliability, or completeness of these data for individual or aggregate use with other data. Original data were compiled from various sources. Spatial information may not meet National Map Accuracy Standards. This information may be updated without notification.



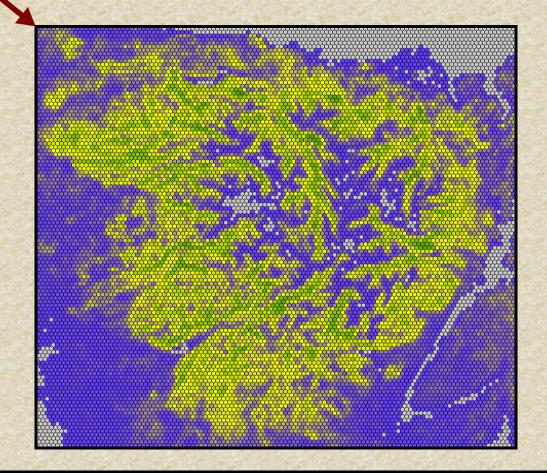


Simulated Spotted Owl Life Cycle



Highest Quality

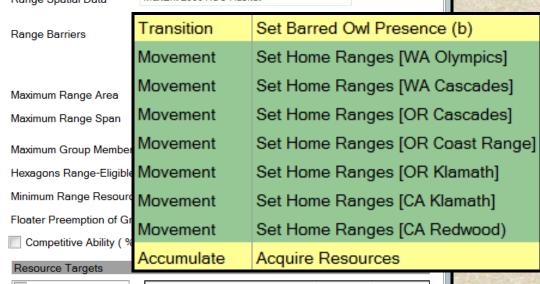
MaxEnt Current Conditions Resource Map



Lowest Quality

Spotted Owl Modeling Regions

Process Varies with Location



	Barred Owl Present
	Demographic Study
✓	Modeling Region
	Resource Class
	Stage Class
	Territory Status

Name	Rank	Target
Not In A Modeling Region	0	0
North Coast Olympics	0	1250
Oregon Coast	0	375
East Cascades South	0	750
East Cascades North	0	1000
West Cascades North	0	1250
West Cascades Central	0	1250
West Cascades South	0	375
Klamath East	0	375
Klamath West	0	375
Inner California Coast Ranges	0	375
Redwood Coast	0	250

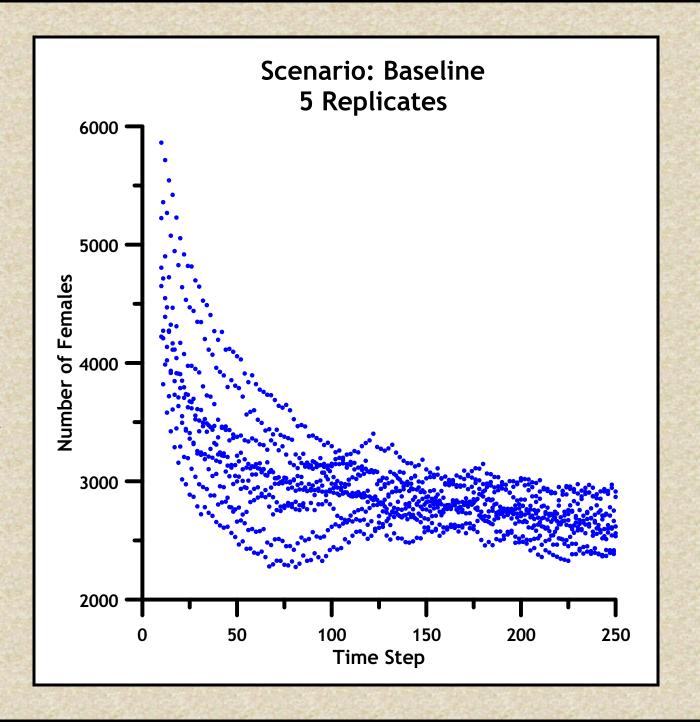
Import

Recover

Close

Results

Total
Population
Size



Results

Population Size by Region

